Rapid Screening to Identify Unusual Thermal Starch Traits from Bulked Corn Kernels

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ABSTRACT

Cereal Chem. 81(4):527-532

Differential scanning calorimetry (DSC) is used routinely to screen for starch thermal properties. In early generations of line development, the established analysis separately evaluates starch extracted from five, single corn kernels. A thermal property trait carried by a recessive gene would appear 25% of the time; thus, if five separate kernels were evaluated, the likelihood of detecting an unusual thermal trait is high. The objective of the current work was to expedite selection by examining five kernels at a time, instead of one, hypothesizing that we would be able to detect different thermal properties in this blend. Corn lines, all from the same

genetic background (ExSeed68 or Oh43), with known thermal functions (amylose-extender, dull, sugary-1, sugary-2, and waxy) were blended with normal starch (control) in ratios of 0:5, 1:4, 2:3, 3:2, 4:1, and 5:0, and analyzed with DSC. The values for each ratio within a mutant type were unique (α < 0.01) for most DSC measurements, especially for gelatinization onset temperature, change in enthalpy of gelatinization, and range of gelatinization. These results support the five-kernel method for rapidly screening large amounts of corn germplasm to identify kernels with unusual starch traits.

Differential scanning calorimetry (DSC) was first utilized by Stevens and Elton (1971) to study starch gelatinization. DSC is an excellent method for researching starch gelatinization because it allows the use of a wide range of starch-to-water ratios, is not limited to temperatures <100°C, and estimates transition enthalpies (Biliaderis et al 1980). Also, DSC requires only a small amount of sample, is easy to operate, and is relatively rapid compared with other methods (Sanders et al 1990; Campbell et al 1995). These factors make it conducive for breeding programs in which large numbers of corn genotypes are screened for desirable starch properties such as low gelatinization onset temperature ($T_{\rm oG}$) or low change in enthalpy of gelatinization ($\Delta H_{\rm G}$).

Variations in DSC measurements have been demonstrated for a variety of maize mutants including amylose-extender (ae), dull (du), sugary-1 (su1), sugary-2 (su2), and waxy (wx) (Inouchi et al 1984; Brockett et al 1988; Ninomya et al 1989; Sanders et al 1990; Inouchi et al 1991; Wang et al 1992; Campbell et al 1995; Perera et al 2001; Tziotis 2001). These particular mutants cause changes from normal corn starch in amylose percentage and phytoglycogen accumulation (Shannon and Garwood 1984). For example, the ae mutation results in starch with 50-70% apparent amylose content (Ikawa et al 1981; Yeh et al 1981; Shannon and Garwood 1984), which may dilute the crystalline regions, thus causing a loss of cooperative melting (Wang et al 1992). The ae mutation also was reported to increase the chain length of amylopectin (Ikawa et al 1981), which would then require a higher temperature to gelatinize (Wang et al 1992). Therefore, the ae starch typically has a broad gelatinization peak that is not complete until up to 120°C, and a high $\Delta H_{\rm G}$ (Brockett et al 1988; Inouchi et al 1991). The du and sul genotypes also are reported to increase apparent amylose percentage (Ikawa et al 1981; Yeh et al 1981) but do not have broad gelatinization peaks typical of ae starch (Inouchi et al 1984; Brockett et al 1988; Wang et al 1992). They both, however, typically possess a lower $\Delta H_{\rm G}$ value and a $T_{\rm oG}$ a few degrees below that of normal starch, which may be a result of slightly lower and less perfect crystallinity in the starch (Inouchi et al 1984). Starch from the su2

Recently, there has been interest in developing corn starches that naturally possess properties similar to those of chemically modified corn starches. The Germplasm Enhancement of Maize (GEM) project has developed and identified exotic by adapted lines that are partially from germplasm foreign to corn races grown in the United States and which may be useful for agronomic, nutritional, and industrial reasons (Pollak 2002). Our laboratory routinely screens corn sources for starch traits that may be useful to the starch industry such as low T_{oG} or low percentage of retrogradation (%R), as well as other criteria (Seetharaman et al 2001). As described earlier, DSC is one of the most rapid methods available for such screening. Earlier methods, however, typically involved extracting starch from single corn kernels such as utilized by Ji et al (2003), for a total of up to 10 kernels from one source. Obanni and BeMiller (1995) described a technique of screening corn through ghost structures, which are the remnants of starch granules after autoclaving small amounts of starch. The greatest value for starch, however, is in the thermal properties. It would be advantageous to expedite the procedure by bulk-extracting starch from a pool of kernels instead of only one kernel, while still being able to recognize the presence of starch with different properties through DSC analysis. Obanni and BeMiller (1997) studied properties of blends of different types of starches such as normal corn, waxy corn, amylose-extender corn, potato, and wheat. They reported that the DSC output did not resemble either of the components in the mixture. However, Liu and Lelievre (1992) reported that the DSC endotherms for blends of wheat and rice starch were the sum of the outputs for each of the components when the starch concentration was <30%. Preliminary data in our laboratory indicated similar results when two starch types were blended together. For example, we used a mixture of normal and su2 starch in different ratios in the DSC pan, which resulted in independent peaks on the DSC, both similar to their respective

The objectives of this study were to investigate the use of DSC as a screening method for detecting unique thermal properties in a blend of two starch types, and to determine whether the starches

genotype also has a higher apparent amylose content than normal starch but gelatinizes at a much lower temperature and $\Delta H_{\rm G}$, which may be a result of the very low percentage of crystallinity and higher amount of short branch-chains of amylopectin in su2 starches than in normal starches (Inouchi et al 1984; Perera et al 2001). The wx genotype causes an elimination of amylose content, unlike the other mutants presented here (Inouchi et al 1984). This mutant results in starch with $\approx 100\%$ amylopectin, the crystalline component of starch, which requires more energy to gelatinize (Inouchi et al 1984).

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gelatinized independently when mixed in different ratios in the DSC pan in excess water. The practical purpose for this work would to be able to detect the presence of a recessive gene affecting starch gelatinization within a segregating corn type. We created model systems with different ratios of mutant (ae, du, su1, su2, and wx) to normal starches to determine the number of kernels containing starch with different functional properties needed in a bulk extraction to be detectable by DSC.

MATERIALS AND METHODS

Materials

Corn (Zea mays L.) lines, all from the same genetic background (ExSeed68), with known thermal functions (amylose-extender 25 [ae], dull 39 [du], sugary-1 [su1], sugary-2 [su2], and waxy 55 [wx]) were obtained from ExSeed Genetics (Ames, IA)

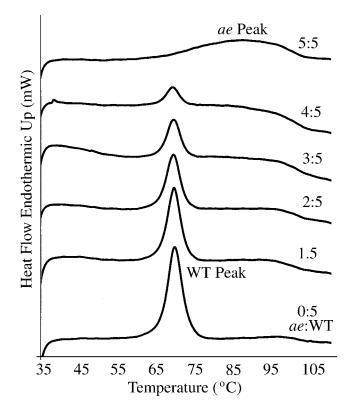


Fig. 1. Differential scanning calorimetry (DSC) thermogram of ratios of *ae* and wild type (WT) corn starch.

along with the wild type of the ExSeed68 background (WT), with normal corn starch. The genes of corn lines from the same background were identical, except for those modified by the genetic mutation. The *ae*, *du*, *wx*, and WT lines were grown and harvested in Ames, IA, during the summer of 1999. The *wx*, *su1*, and a second WT control were grown in the summer of 2000. Another *su2* line in the Oh43 background, Oh43*su2*, was obtained from the USDA-ARS (Ames, IA) along with the Oh43 parent line. These lines were grown and harvested in Ames, IA, during the summer of 1989. All ears were harvested at physiological maturity and dried at 35°C until the moisture content reached 12%. All seeds were stored at 4°C and 45% rh until analyzed.

Starch Extraction

Corn starch was extracted as bulked five-kernel samples from each line, according to the method of White et al (1990) with modifications by Krieger et al (1997) and Ji et al (2004). Each starch type was extracted in triplicate with replicates analyzed separately with DSC.

Differential Scanning Calorimetry

Gelatinization characteristics of starch samples were determined by using differential scanning calorimetry (DSC) (Perkin Elmer DSC 7, Norwalk, CT) equipped with thermal analysis software (Perkin Elmer).

Samples were weighed on the same balance (Mettler AE 104, Toledo, OH). The starches with known thermal properties (ae, du, wx, su1, su2, Oh43su2) were blended in an aluminum DSC pan with their respective background starch, either WT or Oh43 grown in the same season as the respective mutant, in ratios of 0:5, 1:4, 2:3, 3:2, 4:1, and 5:0 (mutant starch [dry weight] to background starch [dry weight]) to give a total starch weight of 3.50 mg. These ratios are reported here as mutant starch ratios 0, 1, 2, 3, 4, and 5, respectively. This plan allowed visualization of the effect of one kernel out of five being unique, two out of five, and so on, up to five out of five kernels. Water was added to the blended starch sample in a water-to-starch ratio of 2:1 and the sample was allowed to equilibrate for at least 30 min before DSC analysis. All samples were analyzed by DSC from 30 to 110°C at 10°C/min. Starch from the ae mutation typically does not fully gelatinize until ≈120°C, but preliminary results showed that unique properties were still detectable up to 110°C. Data parameters collected from the computer included gelatinization onset temperature (T_{oG}) , gelatinization peak temperature (T_{pG}) , gelatinization conclusion temperature (T_{cG}) , and change in enthalpy of gelatinization (ΔH_G) . The $\Delta H_{\rm G}$ was calculated on a starch dry weight basis. Also calculated were the range of gelatinization (R_G) $(T_{cG} \text{ minus } T_{oG})$ and peak height index (PHI) [ΔH_G (dry basis)/(1/2 × R_G)].

TABLE I
Thermal Properties^a of Corn Starch Blends with Ratios of *ae* Starch and Wild Type (WT) Starch

								· -				
	Peak 1a ^b						Peak 1b ^b					
Ratioc	T_{oG} (°C)	<i>T</i> _{pG} (°C)	<i>T</i> _{cG} (°C)	R _G (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})	<i>T</i> _{oG} (°C)	<i>T</i> _{pG} (°C)	<i>T</i> _{cG} (°C)	R _G (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})
0	65.1ab	69.1a	73.1a	8.0a	11.6a	2.9a	87.1a	95.9a	102.4ab	15.3d	1.6d	0.21c
1	65.0a	69.1a	72.8b	7.7ab	8.9b	2.3b	83.7a	96.0a	102.7ab	19.0cd	1.9cd	0.20c
2	65.0a	69.1a	72.6bc	7.6b	6.5c	1.7c	84.0a	95.9a	103.2a	19.2cd	2.5c	0.27bc
3	65.0a	69.1a	72.5c	7.4b	4.3d	1.2d	82.8a	96.0a	103.1a	20.2c	3.4b	0.34b
4	65.4b	69.1a	72.0d	6.6c	1.7e	0.5e	76.5b	93.8b	102.6ab	26.1b	4.2b	0.32b
5							68.2c	84.9c	101.9b	33.8a	11.3a	0.67a
r^{d}	0.17	0.04	-0.87	-0.65	-0.99	-0.99	-0.86	-0.70	-0.19	0.86	0.82	0.82
P	0.3746	0.838	< 0.0001	< 0.0001	< 0.0001	< 0.0001	< 0.0001	< 0.0001	0.27	< 0.0001	< 0.0001	< 0.0001

^a T_{oG} , gelatinization onset temperature; T_{pG} , gelatinization peak temperature; T_{cG} , gelatinization conclusion temperature; R_{G} , range of gelatinization temperature; ΔH_{G} , change in enthalpy of gelatinization; PHI, peak height index ($\Delta H/1/2R_{\text{G}}$). Values in a column with different letters are significantly different (a = 0.01).

b Peak 1a is the WT peak (normal). Peak 1b is the ae peak (unusual starch), except for ratio 0. Peak 1b shows the amylose-lipid complex thermal properties.

^c Each ratio is a proportion of unusual starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

^d Pearson's correlation coefficient of means within a column; P value tests Ho: r = 0.

Statistical Analysis

Calculations were performed with SAS statistical software (v. 8.02) (SAS Institute, Cary, NC). Analysis of variance (ANOVA) was used to test the hypothesis that means were not different within each DSC parameter for each ratio within a mutant type. Tukey's multiple range test was used to test for differences between ratios within a DSC parameter ($\alpha = 0.01$). Pearson correlation coefficients also were calculated for each DSC parameter within each mutant starch.

RESULTS AND DISCUSSION

The model systems were based on a total of five kernels because a recessive trait occurs 25% of the time in a heterozygous population (Poehlman 1959). Therefore, bulk-extracting five kernels greatly increased the probability of detecting a recessive trait. Each unique starch (ae, du, su1, wx, su2, and Oh43su2) imparted properties in the starch mixtures in their own specific ways, such as lowering $T_{\rm oG}$, decreasing $\Delta H_{\rm G}$, or increasing $T_{\rm cG}$.

ae

Because ae starch typically gelatinizes as a broad DSC peak that has only minimal impact in the gelatinization range of amylopectin (Fig. 1), when measured alone as ratio 5, its T_{cG} and R_{G} are very high (Table I). The presence of ae starch was visible in ratios 1 through 4 as a long, broad peak following the peak of the WT (Fig. 1), similar to the typical amylose-lipid peak and created some significant differences in the DSC values. Because the respective peaks for the ae and WT were different, two peaks were measured, peak 1a for WT and peak 1b for ae. For ratio 0, which contained 100% WT starch, the amylose-lipid complex was measured as peak 1b to demonstrate the differences in parameters with the presence of ae starch. The ae starch had greater T_{oG} and greater T_{cG} , but lower ΔH_{G} , than WT starch, agreeing with previous work (Brockett et al 1988). When ae starch was blended with WT starch, peak 1a (WT starch) decreased in ΔH_G (r = -0.99, P <0.0001) with increased amounts of ae starch. The R_G of peak 1a also decreased (r = -0.65, P < 0.0001) from ratios 0 to 4 and the $T_{\rm cG}$ of peak 1a decreased from ratio 0 to 4 (r = -0.87, P <0.0001). The PHI of peak 1a also decreased as the ratio of ae to WT starch increased from ratios 0 to 4 (r = -0.99, P < 0.0001). The T_{oG} of peak 1b decreased with increased amounts of *ae* starch (r = -0.86, P < 0.0001). The T_{oG} of peak 1b, however, was probably lower than measured in the presence of WT starch because

TABLE II Thermal Properties^a of Corn Starch Blends with Ratios of du Starch and Wild Type (WT) Starch

Ratiob	<i>T</i> _{oG} (°C)	<i>T</i> _{pG} (°C)	<i>T</i> _{cG} (°C)	$R_{\mathbf{G}}$ (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})
0	64.8a	69.0a	73.2a	8.4c	11.7a	2.80a
1	64.3ab	68.8a	73.2a	8.9bc	11.8a	2.69a
2	63.7bc	68.8a	73.3a	9.6ab	10.9ab	2.28b
3	63.3cd	68.4b	73.3a	10.0a	10.4ab	2.09bc
4	62.9d	68.2b	73.2a	10.3a	10.7ab	2.08bc
5	62.8d	67.7c	73.1a	10.3a	9.8b	1.90c
r^{c}	-0.89	-0.86	-0.07	0.83	-0.69	-0.83
P	< 0.0001	< 0.0001	0.7021	< 0.0001	< 0.0001	< 0.0001

^a $T_{\rm oG}$, gelatinization onset temperature; $T_{\rm pG}$, gelatinization peak temperature; $T_{\rm cG}$, gelatinization conclusion temperature; $R_{\rm G}$, range of gelatinization temperature; $\Delta H_{\rm G}$, change in enthalpy of gelatinization; PHI, peak height index ($\Delta H/_{1/2}R_{\rm G}$). Values in a column with different letters are significantly different (a=0.01).

its onset occurred within the range of the WT starch. The $R_{\rm G}$ of peak 1b increased with higher amounts of ae starch (r=0.86, P<0.0001). The $\Delta H_{\rm G}$ of peak 1b also increased with higher amounts of ae starch (r=0.82, P<0.0001) along with the PHI of peak 1b (r=0.82, P<0.0001).

These results indicated that the low crystallinity of the ae starch (Wang et al 1992) altered thermal properties such as increasing the $R_{\rm G}$, and these differences were detectable by DSC. The detection limit in the ae model system was a 1:5 kernel ratio for $\Delta H_{\rm G}$ and PHI of peak 1a, a 2:5 kernel ratio for $T_{\rm cG}$ of peak 1a and $\Delta H_{\rm G}$ of peak 1b, a 3:5 kernel ratio for $R_{\rm G}$ and PHI of peak 1b, and a 4:5 kernel ratio for $R_{\rm G}$ of peak 1a. In practical terms, the detection limit is the ratio of kernels needed to create a significant difference in a specific DSC parameter. The presence of ae starch was visible on a DSC curve and measurable by $\Delta H_{\rm G}$ and PHI for the WT peak at ratio 1.

These results suggested that an *ae*-type starch could be detected in a blend of bulked-extracted starch from five kernels, even if only one unique kernel was present.

The DSC operator could then return to the original line source of the bulked sample and extract using single-kernel methods to locate the kernel source of the *ae*-like starch.

TABLE III Thermal Properties of Corn Starch Blends with Ratios of suI Starch and Wild Type (WT) Starch

Ratiob	<i>T</i> _{oG} (°C)	<i>T</i> _{pG} (°C)	<i>T</i> _{cG} (°C)	<i>R</i> _G (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})
0	66.0a	69.7a	73.7c	7.7c	12.0a	3.12a
1	65.3b	69.8a	73.9bc	8.6b	11.8a	2.74b
2	63.7c	70.0a	74.4a	10.6a	11.9b	2.09c
3	63.0d	69.9a	74.3ab	11.3a	10.9bc	1.93cd
4	63.1d	68.3b	74.3ab	11.2a	10.4cd	1.85d
5	63.1d	67.3c	73.8c	10.7a	9.9d	1.86d
r^{c}	-0.88	-0.79	0.13	0.78	-0.81	-0.89
P	< 0.0001	< 0.0001	0.4282	< 0.0001	< 0.0001	< 0.0001

^a $T_{\rm oG}$, gelatinization onset temperature; $T_{\rm pG}$, gelatinization peak temperature; $T_{\rm cG}$, gelatinization conclusion temperature; $R_{\rm G}$, range of gelatinization temperature; $\Delta H_{\rm G}$, change in enthalpy of gelatinization; PHI, peak height index ($\Delta H/1/2R_{\rm G}$). Values in a column with different letters are significantly different (a = 0.01).

TABLE IV
Thermal Properties^a of Corn Starch Blends
with Ratios of wx Starch and Wild Type (WT) Starch

0 66.1a 69.9c 73.9e 7.8f 12.0b 3.06a 1 66.0ab 69.9c 74.3e 8.3e 12.3b 2.98ab 2 65.8bc 70.0c 74.9d 9.1d 12.8ab 2.82b 3 65.5cd 70.1c 75.7c 10.2c 12.9ab 2.53c 4 65.4de 70.4b 76.4b 11.0b 13.1ab 2.39cc 5 65.2e 70.9a 77.4a 12.2a 13.6a 2.22d re -0.78 0.64 0.93 0.98 0.74 0.95							
1 66.0ab 69.9c 74.3e 8.3e 12.3b 2.98at 2 65.8bc 70.0c 74.9d 9.1d 12.8ab 2.82b 3 65.5cd 70.1c 75.7c 10.2c 12.9ab 2.53c 4 65.4de 70.4b 76.4b 11.0b 13.1ab 2.39cc 5 65.2e 70.9a 77.4a 12.2a 13.6a 2.22d re -0.78 0.64 0.93 0.98 0.74 0.95	Ratiob	<i>T</i> _{oG} (°C)	<i>T</i> _{pG} (°C)	<i>T</i> _{cG} (°C)	<i>R</i> _G (°C)	•	PHI (Δ <i>H</i> /°C)
2 65.8bc 70.0c 74.9d 9.1d 12.8ab 2.82b 3 65.5cd 70.1c 75.7c 10.2c 12.9ab 2.53c 4 65.4de 70.4b 76.4b 11.0b 13.1ab 2.39cc 5 65.2e 70.9a 77.4a 12.2a 13.6a 2.22d rc -0.78 0.64 0.93 0.98 0.74 0.95	0	66.1a	69.9c	73.9e	7.8f	12.0b	3.06a
3 65.5cd 70.1c 75.7c 10.2c 12.9ab 2.53c 4 65.4de 70.4b 76.4b 11.0b 13.1ab 2.39cc 5 65.2e 70.9a 77.4a 12.2a 13.6a 2.22d re -0.78 0.64 0.93 0.98 0.74 0.95	1	66.0ab	69.9c	74.3e	8.3e	12.3b	2.98ab
4 65.4de 70.4b 76.4b 11.0b 13.1ab 2.39cc 5 65.2e 70.9a 77.4a 12.2a 13.6a 2.22d rc -0.78 0.64 0.93 0.98 0.74 0.95	2	65.8bc	70.0c	74.9d	9.1d	12.8ab	2.82b
5 65.2e 70.9a 77.4a 12.2a 13.6a 2.22d rc -0.78 0.64 0.93 0.98 0.74 0.95	3	65.5cd	70.1c	75.7c	10.2c	12.9ab	2.53c
r^{c} -0.78 0.64 0.93 0.98 0.74 0.95	4	65.4de	70.4b	76.4b	11.0b	13.1ab	2.39cd
	5	65.2e	70.9a	77.4a	12.2a	13.6a	2.22d
P <0.0001 <0.0001 <0.0001 <0.0001 <0.0001 <0.0001	r^{c}	-0.78	0.64	0.93	0.98	0.74	0.95
	P	< 0.0001	< 0.0001	< 0.0001	< 0.0001	< 0.0001	< 0.0001

^a $T_{\rm oG}$, gelatinization onset temperature; $T_{\rm pG}$, gelatinization peak temperature; $T_{\rm cG}$, gelatinization conclusion temperature; $R_{\rm G}$, range of gelatinization temperature; $\Delta H_{\rm G}$, change in enthalpy of gelatinization; PHI, peak height index ($\Delta H_{\rm G}/R_{\rm G}$). Values in a column with different letters are significantly different (a=0.01).

^b Each ratio is a proportion of *du* starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

^c Pearson's correlation coefficient of means within a column; P value tests Ho: r = 0.

^b Each ratio is a proportion of *su1* starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

^c Pearson's correlation coefficient of means within a column; P value tests are Ho: r = 0.

^b Each ratio is a proportion of wx starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

^c Pearson's correlation coefficient of means within a column; P value tests Ho: r=0.

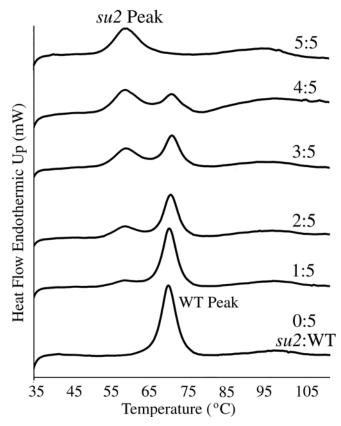


Fig. 2. Differential scanning calorimetry (DSC) thermogram of ratios of *su2* and wild type (WT) corn starch.

du

The $T_{\rm oG}$ and $\Delta H_{\rm G}$ of 100% du starch (ratio 5) were lower than these values for 100% WT starch (ratio 0), which agreed with previous work (Brockett et al 1988; Inouchi et al 1991; Wang et al 1992) (Table II). The $T_{\rm oG}$ decreased with increased amounts of du starch (r=-0.89, P<0.0001). The decrease in $T_{\rm oG}$ was accompanied by an increase in $R_{\rm G}$ (r=0.83, P<0.0001). The PHI also decreased with increased ratios (r=-0.83, P<0.0001). The $\Delta H_{\rm G}$, however, did not become significantly different from the WT starch until ratio 5 (100% du starch), but the value did decrease with an increase in ratio of du starch (r=-0.69, P<0.0001). Inouchi et al (1984) reported that du starch was less crystalline than normal starch, thus causing changes in thermal DSC properties.

In the current work, all parameters decreased, except for $R_{\rm G}$, with increased amounts of the du starch. The detection limit for du starch was a 2:5 kernel ratio with respect to $T_{\rm oG}$, $R_{\rm G}$, and PHI and 3:5 for $T_{\rm pG}$. These results suggest that the presence of a du type starch could be detected in a bulk extraction of five kernels, when only two kernels had the recessive trait, by identifying a lower $T_{\rm oG}$ and $\Delta H_{\rm G}$ than normal.

As noted previously for the *ae*-type starch, the investigator could then return to the original corn line and re-extract by single-kernel methods to identify additional kernels with this less crystalline starch.

sul

The suI starch had lower T_{oG} , T_{pG} , ΔH_{G} , and wider R_{G} than the WT starch, which was similar to previous reports of suI starch in various backgrounds (Brockett et al 1988; Inouchi et al 1991; Wang et al 1992) (Table III). T_{oG} decreased with increased amounts of suI starch (r = -0.88, P < 0.0001). There were few differences, however, among blends at higher ratios of suI starch (3, 4, and 5). The decrease in T_{oG} was accompanied by an increase in R_{G} (r = -0.88).

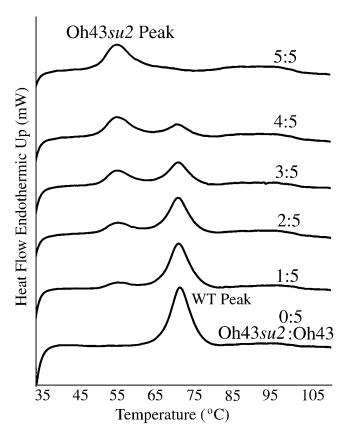


Fig. 3. Differential scanning calorimetry (DSC) thermogram of ratios of Oh43*su2* and Oh43 corn starch.

0.78, P < 0.0001). Similarly, the differences were not significant at higher ratios (3, 4, and 5). The $\Delta H_{\rm G}$ decreased significantly with increased increments of su1 starch (r = -0.81, P < 0.0001). The PHI also decreased significantly (r = -0.89, P < 0.0001). Overall, these results indicated that small amounts of su1 starch significantly affected the DSC results. The su1 starch was lower in crystallinity than normal starch, which resulted in lower $T_{\rm oG}$ and $\Delta H_{\rm G}$ (Inouchi et al 1984).

These differences were significant in the model systems at a 1:5 kernel ratio for $T_{\rm oG}$, $R_{\rm G}$, and PHI, a 2:5 kernel ratio for $\Delta H_{\rm G}$, and a 4:5 kernel ratio for $T_{\rm pG}$, the latter which decreased significantly with the addition of suI starch (r = -0.79, P < 0.0001).

wx

The thermal properties of the wx starch had lower T_{oG} , greater T_{cG} , wider R_{G} , greater ΔH_{G} , and lower PHI than the WT starch (Table IV). These results agreed with previous findings of wx starch in another background (Inouchi et al 1984) and were a consequence of the higher crystallinity of the wx starch (Inouchi et al 1984). In the current study, the $T_{\rm oG}$ tended to decrease (r = -0.78, P < 0.0001), whereas the T_{cG} increased (r = 0.93, P <0.0001), causing a wider R_G (r = 0.98, P < 0.0001) with greater ratios of wx starch. The $\Delta H_{\rm G}$ also increased with increased amounts of wx starch but only became significantly different from the WT starch at ratio 5 (100% wx starch) (r = 0.74, P < 0.0001). The PHI also decreased significantly as the ratio of wx starch increased (r = -0.95, P < 0.0001). As the ratio of wx starch to WT increased, the crystallinity of the starch blend increased (Inouchi et al 1984), causing a rise in $\Delta H_{\rm G}$ and $R_{\rm G}$, which resulted in a lower PHI.

The differences were significant at a 1:5 kernel ratio for $R_{\rm G}$, a 2:5 kernel ratio for $T_{\rm oG}$, $T_{\rm cG}$, and PHI and a 4:5 kernel ratio for $T_{\rm pG}$, the latter which increased with the addition of wx starch (r = 0.64, P < 0.0001).

TABLE V
Thermal Properties^a of Corn Starch Blends with Ratios of su2 Starch and Wild Type (WT) Starch

	Peak 1a ^b							Peak 1b ^b				
Ratioc	<i>T</i> _{oG} (°C)	T_{pG} (°C)	T _{cG} (°C)	R_{G} (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})	<i>T</i> _{oG} (°C)	T_{pG} (°C)	<i>T</i> _{cG} (°C)	R _G (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})
0							65.0a	69.1a	73.3a	8.3a	11.7a	2.82a
1	53.7a	57.4a	59.9a	6.2a	0.4a	0.11a	65.2ab	69.3ab	73.3a	8.1ab	7.9b	1.98b
2	52.9b	57.3a	61.2b	8.3b	1.1b	0.27b	65.4ab	69.4a-c	73.1a	7.7bc	5.5c	0.94c
3	52.6b	57.5a	61.9c	9.4c	2.2c	0.47c	65.8b	69.6bc	73.2a	7.4c	3.2d	0.58d
4	52.5b	57.5a	62.6d	10.2d	3.4d	0.66d	66.6c	69.8c	73.2a	6.6d	1.6e	0.30e
5	51.8c	57.5a	64.0e	12.3e	6.8e	1.10e						
r^{d}	-0.57	0.16	0.87	0.96	0.94	0.97	0.55	0.55	-0.13	-0.66	-0.98	-0.94
P	0.0011	0.3858	< 0.0001	< 0.0001	< 0.0001	< 0.0001	0.0015	0.0014	0.5041	< 0.0001	< 0.0001	< 0.0001

^a T_{oG} , gelatinization onset temperature; T_{pG} , gelatinization peak temperature; T_{cG} , gelatinization conclusion temperature; R_{G} , range of gelatinization temperature; ΔH_{G} , change in enthalpy of gelatinization; PHI, peak height index ($\Delta H/1/2R_{\text{G}}$). Values in a column with different letters are significantly different (a = 0.01).

u2 and Oh43su2

Because the $T_{\rm oG}$ and $T_{\rm cG}$ of the su2 and Oh43su2 starches were much lower than those of their background starches, two separate gelatinization peaks were formed when they were blended (Figs. 2 and 3). The first peak (1a) was caused by the presence of either su2 or Oh43su2 starch and the second peak (1b) was contributed by the background starch, either WT or Oh43.

With greater proportions of su2 starch, the $\Delta H_{\rm G}$ of peak 1a increased (r=0.94, P<0.0001), whereas the $\Delta H_{\rm G}$ of peak 1b decreased (r=-0.98, P<0.0001) (Table V). The $R_{\rm G}$ of peak 1a increased (r=0.96, P<0.0001), whereas the $R_{\rm G}$ of peak 1b decreased (r=-0.66, P<0.0001) with greater proportions of su2 starch.

This pattern resulted in increased PHI of peak 1a (r=0.97, P<0.0001) and decreased PHI of peak 1b (r=-0.94, P<0.0001). Total $\Delta H_{\rm G}$ of the su2 model system (Table VI) decreased with an increased proportion of su2 starch (r=-0.94, P<0.0001). The detection limit was a 1:5 kernel ratio for $\Delta H_{\rm G}$ and PHI of peak 1b, and for $R_{\rm G}$ and PHI of the total starch sample. A 2:5 kernel ratio was needed to detect significant differences in $T_{\rm oG}$, $T_{\rm cG}$, $T_{\rm cG}$, $T_{\rm cG}$, $T_{\rm cG}$, and PHI of peak 1a, and $T_{\rm cG}$ of peak 1b. To detect differences in $T_{\rm cG}$ and $T_{\rm cG}$ of peak 1b, and $T_{\rm cG}$ of the total starch sample, a kernel ratio of 3:5 was required.

Similar to the su2 results, the $\Delta H_{\rm G}$ of peak 1a increased with greater proportions of Oh43su2 starch, (r = 0.94, P < 0.0001), whereas the $\Delta H_{\rm G}$ of peak 1b decreased (r = -0.99, P < 0.0001) (Table VII). The R_G of peak 1a also increased (r = 0.89, P <0.0001), whereas the R_G of peak 1b decreased (r = -0.79, P <0.0001) with increased proportions of the Oh43su2 starch. This pattern caused the PHI of peak 1a to also increase (r = 0.95, P <0.0001) and the PHI of peak 1b to decrease (r = -0.98, P <0.0001). The total $\Delta H_{\rm G}$ (Table VIII) decreased significantly as the ratio of the Oh43su2 starch increased (r = -0.91, P < 0.0001). These differences were significant at a 1:5 kernel ratio for $\Delta H_{\rm G}$ and PHI of peak 1b, and for R_G and PHI of the total, combined peaks. A 2:5 kernel ratio was needed to identify differences in the T_{cG} , R_G , ΔH_G , and PHI of peak 1a, and ΔH_G of the total combined peaks. A 3:5 kernel ratio was needed to achieve significant differences in T_{cG} and R_{G} for peak 1b, and a 4:5 kernel ratio was needed to identify differences in T_{oG} of peaks 1a and 1b.

Overall, detection of unique properties for the Oh43su2 and su2 model systems was visible at a 1:5 kernel ratio because a separate peak occurred that was a result of the su2 starch (Figs. 2 and 3). These ratio studies demonstrated how the thermal properties of a starch mixture could be affected by having just 20% of a starch that is different from the background starch. The su2 and Oh43su2 starches were lower in crystallinity than normal starches (Inouchi et al 1984; Perera et al 2001) and therefore resulted in lower T_{oG}

TABLE VI
Total Thermal Properties^a of Corn Starch Blends
with Ratios of su2 Starch and Wild Type (WT) Starch

	• • • • • • • • • • • • • • • • • • • •									
	Total ^b									
Ratioc	$\Delta H_{\rm G}$ (J/g)	<i>R</i> _G (°C)	PHI ($\Delta H/_{1/2R_G}$)							
0	11.7a	8.3a	2.82a							
1	10.6a	19.6b	1.09b							
2	10.2ab	20.2bc	1.01bc							
3	8.9bc	20.6c	0.86cd							
4	7.7cd	20.7c	0.74d							
5	6.8d	12.3d	1.10b							
r^{d}	-0.94	0.24	-0.66							
P	< 0.0001	0.1673	< 0.0001							

^a $T_{\rm oG}$, gelatinization onset temperature; $T_{\rm pG}$, gelatinization peak temperature; $T_{\rm cG}$, gelatinization conclusion temperature; $R_{\rm G}$, range of gelatinization temperature; $\Delta H_{\rm G}$, change in enthalpy of gelatinization; PHI, peak height index ($\Delta H/_{1/2}R_{\rm G}$). Values in a column with different letters are significantly different (a=0.01).

and ΔH_G , which then produced a peak that was entirely separate from that created from the gelatinization peaks of the WT or Oh43 starch.

CONCLUSIONS

This study demonstrated that a five-kernel bulk extraction might be used to screen corn starch for the presence of only one in five kernels with different thermal properties. If one kernel out of five were different from the normal starch, it was detected by DSC, as demonstrated with model systems created by blending different ratios of mutant starches previously shown to possess thermal properties different from those found in normal corn (ae, du, su1, wx, su2, and Oh43su2). A five-kernel bulk extraction was very helpful in expediting the screening process because it reduced the extraction time and DSC analyses by a factor of five. This method can be applied to screening unknown germplasm for starch thermal properties that are different from the properties of normal corn starch. The study also demonstrated that starch blends do impart the thermal properties of the independent starch components, suggesting conclusions that differed from those of Obanni and BeMiller (1997) and agreeing with those of Liu and Lelievre (1992) because the starch blends in our experiments were in excess

^b Peak 1a is the Oh43su2 peak (unusual starch). Peak 1b is the WT peak (normal starch) peak.

^c Each ratio is a proportion of unusual starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

d Pearson's correlation coefficient of means within a column; P value tests Ho: r = 0.

b Total is the result of the analysis from the start of Peak 1a to the end of Peak 1b.

^c Each ratio is a proportion of unusual starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

^d Pearson's correlation coefficient of means within a column; P value tests Ho: r = 0.

TABLE VII
Thermal Properties^a of Corn Starch Blends with Ratios of Oh43su2 Starch and Wild Type (WT) Starch

	Peak 1a ^b							Peak 1b ^b				
Ratioc	<i>T</i> _{oG} (°C)	<i>T</i> _{pG} (°C)	T _{cG} (°C)	$R_{\rm G}$ (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})	<i>T</i> _{oG} (°C)	<i>T</i> _{pG} (°C)	T _{cG} (°C)	R _G (°C)	$\Delta H_{\rm G}$ (J/g)	PHI (Δ <i>H</i> / _{1/2RG})
0							67.2a	71.8a	77.0a	9.8a	11.9a	2.41a
1	51.4a	55.7a	59.8a	8.5a	0.7a	0.16a	67.1a	71.7a	76.8ab	9.7a	9.0b	1.87b
2	51.3a	56.1a	60.9b	9.5b	1.6b	0.35b	67.4a	71.9a	76.9ab	9.5ab	6.2c	1.30c
3	50.9ab	55.9a	60.9b	10.0b	2.5c	0.51c	67.2a	71.6a	76.3bc	9.1b	3.8d	0.83d
4	50.8b	56.0a	61.8c	11.0c	3.9d	0.70d	68.2b	71.7a	75.9c	7.8c	1.5e	0.38e
5	50.6b	56.3a	62.9d	12.3d	7.5e	1.22e						
$r^{ m d}$	-0.33	0.17	0.83	0.89	0.94	0.95	0.44	-0.11	-0.70	-0.79	-0.99	-0.98
P	0.0724	0.3687	< 0.0001	< 0.0001	< 0.0001	< 0.0001	0.0147	0.5764	< 0.0001	< 0.0001	< 0.0001	< 0.0001

^a T_{oG} , gelatinization onset temperature; T_{pG} , gelatinization peak temperature; T_{cG} , gelatinization conclusion temperature; R_{G} , range of gelatinization temperature; ΔH_{G} , change in enthalpy of gelatinization; PHI, peak height index ($\Delta H/1/2R_{\text{G}}$). Values in a column with different letters are significantly different (a = 0.01).

TABLE VIII
Total Thermal Properties^a of Corn Starch Blends with Ratios of Oh43su2 Starch and Wild Type (WT) Starch

	Total ^b									
Ratioc	ΔH_{G} (J/g)	R_{G} (°C)	PHI (Δ <i>H</i> / _{1/2RG})							
0	11.5a	9.8a	2.35a							
1	11.5a	25.4b	0.90b							
2	9.6b	25.5b	0.76bc							
3	8.2c	25.4b	0.65c							
4	7.3c	25.1b	0.59c							
5	7.5c	12.3c	1.22d							
r^{d}	-0.91	0.08	-0.53							
P	< 0.0001	0.6421	0.0008							

^a $T_{\rm oG}$, gelatinization onset temperature; $T_{\rm pG}$, gelatinization peak temperature; $T_{\rm cG}$, gelatinization conclusion temperature; $R_{\rm G}$, range of gelatinization temperature; $\Delta H_{\rm G}$, change in enthalpy of gelatinization; PHI, peak height index ($\Delta H/_{1/2}R_{\rm G}$). Values in a column with different letters are significantly different (a=0.01).

water. In future work, it would be useful to study the impact of these starch blends on pasting properties of the mixtures, which could lend further insight into the thermal interactions of starch types.

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[Received December 9, 2003. Accepted February 25, 2004.]

^b Peak 1a is the Oh43su2 peak (unusual starch). Peak 1b is the WT peak (normal starch) peak.

^c Each ratio is a proportion of unusual starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

^d Pearson's correlation coefficient of means within a column; P value tests Ho: r = 0.

b Total is the result of the analysis from the start of Peak 1a to the end of Peak

^c Each ratio is a proportion of unusual starch to normal starch (ratio 0 = 0 parts unusual starch to 5 parts normal starch).

^d Pearson's correlation coefficient of means within a column; P value tests Ho: r = 0.